

RAW SEQUENCE LISTING

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Application Serial Number: 10/637,710
Source: TELEO
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PATENT APPLICATION: US/10/637,710

DATE: 03/18/2005
TIME: 17:27:17

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Output Set: N:\CRF4\03182005\J637710.raw

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3 <110> APPLICANT: Panda, Satchidananda
4      Hogenesch, John B.
5      Provincio, Ignacio
6      Kay, Steve A.
7      IRM LLC
8      Uniformed Services University of the Health Sciences
10 <120> TITLE OF INVENTION: Methods for Treating Circadian Rhythm Phase
11      Disturbances
13 <130> FILE REFERENCE: 021288-001020US
15 <140> CURRENT APPLICATION NUMBER: US 10/637,710
16 <141> CURRENT FILING DATE: 2003-08-08
18 <150> PRIOR APPLICATION NUMBER: US 60/402,570
19 <151> PRIOR FILING DATE: 2002-08-08
21 <150> PRIOR APPLICATION NUMBER: US 60/482,384
22 <151> PRIOR FILING DATE: 2003-06-25
24 <160> NUMBER OF SEQ ID NOS: 12
26 <170> SOFTWARE: PatentIn Ver. 2.1
28 <210> SEQ ID NO: 1
29 <211> LENGTH: 2137
30 <212> TYPE: DNA
31 <213> ORGANISM: Mus sp.
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34 <223> OTHER INFORMATION: mouse melanopsin cDNA
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39 agccggaagt ctggggaccg atccctgatc tttccatggc cttagctcct ctgagagcct 180
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41 cacaaccagt cctgccctgc aaggcatttg gaacggcact cagaacgtct ccgtaaagagc 300
42 ccagcttctc tctgttagcc ccacgacatc tgcacatcag gctgctgcct gggccccctt 360
43 ccccacagtc gatgtcccag accatgctca ctatacccta ggcacggtga tcctgctgg 420
44 gggactcaca gggatgtgg gcaatctgac ggtcatctac accttctgca ggaacagagg 480
45 cctgcggaca ccagcaaaca ttttcatcat caacctcgca gtcagcgact tcctcatgtc 540
46 agtcaactcag gccccggctc tctttgccag cagcctctac aagaagtgc tctttggga 600
47 gacaggttgc gagttctatg ctttctgccc ggctgtctt ggcacatcact ccatgatcac 660
48 cctgacagcc atagccatgg accgctatct ggtgatcaca cgtccactgg ccaccatcgg 720
49 caggggatcc aaaagacgaa cggcactcgt cctgttaggc gtctggctt atgccttggc 780
50 ctggagtctg ccaccttct ttgggtggag tgcctacgtg cccgaggggc tgctgacatc 840
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52 ctgtttgtc ttcttctcc ccctgtcat catcatctt tgctacatct tcacatctcag 960
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 58 ccagcacctg cttgccttgg ggtgttctt cggttatca ggccagcga gccaccctc 1320
 59 cctcagctac cgctctaccc accgctccac attgagcagc cagtccttag acctcagctg 1380
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 62 tcagaaccta gaagatggag aactcaaggc ctctccagc ccccaaggta agagatctaa 1560
 63 gactcccaag gtgcctggac ccagtacccctt ccgcctatg aaaggacagg gagccaggcc 1620
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 68 tataacttat gtgccttga agatatgtgg cctacacgcg agaacaactc atgcgtgtgt 1920
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 70 acagagtgtg atgggttca cctctctgctc cgggtttga tgctgggcaa acacgggaa 2040
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76 <211> LENGTH: 521

77 <212> TYPE: PRT

78 <213> ORGANISM: Mus sp.

80 <220> FEATURE:

81 <223> OTHER INFORMATION: mouse melanopsin

83 <400> SEQUENCE: 2

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88						20					25					30
90	Gln	Asn	Val	Ser	Val	Arg	Ala	Gln	Leu	Leu	Ser	Val	Ser	Pro	Thr	Thr
91						35					40					45
93	Ser	Ala	His	Gln	Ala	Ala	Ala	Trp	Val	Pro	Phe	Pro	Thr	Val	Asp	Val
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96	Pro	Asp	His	Ala	His	Tyr	Thr	Leu	Gly	Thr	Val	Ile	Leu	Leu	Val	Gly
97	65					70					75					80
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100						85					90					95
102	Asn	Arg	Gly	Leu	Arg	Thr	Pro	Ala	Asn	Met	Phe	Ile	Ile	Asn	Leu	Ala
103						100					105					110
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108	Ser	Ser	Leu	Tyr	Lys	Lys	Trp	Leu	Phe	Gly	Glu	Thr	Gly	Cys	Glu	Phe
109						130					135					140
111	Tyr	Ala	Phe	Cys	Gly	Ala	Val	Phe	Gly	Ile	Thr	Ser	Met	Ile	Thr	Leu
112	145						145					150				155
114	Thr	Ala	Ile	Ala	Met	Asp	Arg	Tyr	Leu	Val	Ile	Thr	Arg	Pro	Leu	Ala
115											165					170
117	Thr	Ile	Gly	Arg	Gly	Ser	Lys	Arg	Arg	Thr	Ala	Leu	Val	Leu	Leu	Gly
118											180					185
120	Val	Trp	Leu	Tyr	Ala	Leu	Ala	Trp	Ser	Leu	Pro	Pro	Phe	Phe	Gly	Trp
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123 Ser Ala Tyr Val Pro Glu Gly Leu Leu Thr Ser Cys Ser Trp Asp Tyr
124      210          215          220
126 Met Thr Phe Thr Pro Gln Val Arg Ala Tyr Thr Met Leu Leu Phe Cys
127      225          230          235          240
129 Phe Val Phe Phe Leu Pro Leu Leu Ile Ile Phe Cys Tyr Ile Phe
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132 Ile Phe Arg Ala Ile Arg Glu Thr Gly Arg Ala Cys Glu Gly Cys Gly
133          260          265          270
135 Glu Ser Pro Leu Arg Gln Arg Arg Gln Trp Gln Arg Leu Gln Ser Glu
136          275          280          285
138 Trp Lys Met Ala Lys Val Ala Leu Ile Val Ile Leu Leu Phe Val Leu
139          290          295          300
141 Ser Trp Ala Pro Tyr Ser Thr Val Ala Leu Val Ala Phe Ala Gly Tyr
142 305          310          315          320
144 Ser His Ile Leu Thr Pro Tyr Met Ser Ser Val Pro Ala Val Ile Ala
145          325          330          335
147 Lys Ala Ser Ala Ile His Asn Pro Ile Ile Tyr Ala Ile Thr His Pro
148          340          345          350
150 Lys Tyr Arg Val Ala Ile Ala Gln His Leu Pro Cys Leu Gly Val Leu
151          355          360          365
153 Leu Gly Val Ser Gly Gln Arg Ser His Pro Ser Leu Ser Tyr Arg Ser
154          370          375          380
156 Thr His Arg Ser Thr Leu Ser Ser Gln Ser Ser Asp Leu Ser Trp Ile
157 385          390          395          400
159 Ser Gly Arg Lys Arg Gln Glu Ser Leu Gly Ser Glu Ser Glu Val Gly
160          405          410          415
162 Trp Thr Asp Thr Glu Thr Thr Ala Ala Trp Gly Ala Ala Gln Gln Ala
163          420          425          430
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172 465          470          475          480
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187 <213> ORGANISM: Homo sapiens
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190 <223> OTHER INFORMATION: human melanopsin cDNA
192 <400> SEQUENCE: 3
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253 50 55 60
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256 65 70 75 80
258 Gly Leu Thr Gly Met Leu Gly Asn Leu Thr Val Ile Tyr Thr Phe Cys
259 85 90 95
261 Arg Ser Arg Ser Leu Arg Thr Pro Ala Asn Met Phe Ile Ile Asn Leu
262 100 105 110
264 Ala Val Ser Asp Phe Leu Met Ser Phe Thr Gln Ala Pro Val Phe Phe
265 115 120 125
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268 130 135 140
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273 Leu Thr Ala Ile Ala Leu Asp Arg Tyr Leu Val Ile Thr Arg Pro Leu
274 165 170 175
276 Ala Thr Phe Gly Val Ala Ser Lys Arg Arg Ala Ala Phe Val Leu Leu
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280 195 200 205
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304 325 330 335
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307 340 345 350
309 Thr His Pro Lys Tyr Arg Val Ala Ile Ala Gln His Leu Pro Cys Leu
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VERIFICATION SUMMARY

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